



## RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:

10/761,435

Source:

IFWO

Date Processed by STIC:

1/30/04

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FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT

MARK SPENCER, TELEPHONE: 703-308-4212; FAX: 703-308-4221

*Effective 12/13/03:* TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER  
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TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkr41note.htm>

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Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/efb/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry directly to (EFFECTIVE 12/01/03):  
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Revised 10/08/03



IFWO

## RAW SEQUENCE LISTING

DATE: 01/30/2004

PATENT APPLICATION: US/10/761,435

TIME: 14:54:06

Input Set : A:\SeqList 1975\_0180003.ST25.txt

Output Set: N:\CRF4\01302004\J761435.raw

3 <110> APPLICANT: Umana, Pablo  
 4 Bruenker, Peter  
 5 Ferrara, Claudia  
 6 Suter, Tobias  
 8 <120> TITLE OF INVENTION: Fusion Constructs and Use of Same to Produce Antibodies with  
 9 Increased Fc Receptor Binding Affinity and Effector Function  
 11 <130> FILE REFERENCE: 1975.0180003  
 C--> 13 <140> CURRENT APPLICATION NUMBER: US/10/761,435  
 C--> 13 <141> CURRENT FILING DATE: 2004-01-22  
 13 <150> PRIOR APPLICATION NUMBER: US 60/441,307  
 14 <151> PRIOR FILING DATE: 2003-01-22  
 16 <150> PRIOR APPLICATION NUMBER: US 60/491,254  
 17 <151> PRIOR FILING DATE: 2003-07-31  
 19 <150> PRIOR APPLICATION NUMBER: US 60/495,142  
 20 <151> PRIOR FILING DATE: 2003-08-15  
 22 <160> NUMBER OF SEQ ID NOS: 20  
 24 <170> SOFTWARE: PatentIn version 3.2

**Does Not Comply**  
**Corrected Diskette Needed**

(p9.1-2, 5-6)

## ERRORED SEQUENCES

718 <210> SEQ ID NO: 17  
 719 <211> LENGTH: 3432 - Found 3435  
 720 <212> TYPE: DNA  
 721 <213> ORGANISM: Homo sapiens  
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 728 ggctccttcc ctcagggcca gctctcaatg ttgcaagaaa aaatagacca tttggagcgt 180  
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E--&gt; 838 atccagttga ggtga

841 &lt;210&gt; SEQ ID NO: 18

842 &lt;211&gt; LENGTH: 1143

843 &lt;212&gt; TYPE: PRT

844 &lt;213&gt; ORGANISM: Homo sapiens

846 &lt;400&gt; SEQUENCE: 18

847 Met Lys Leu Ser Arg Gln Phe Thr Val Phe Gly Ser Ala Ile Phe Cys

848 1 5 10 15

851 Val Val Ile Phe Ser Leu Tyr Leu Met Leu Asp Arg Gly His Leu Asp

3435

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855 Tyr Pro Arg Asn Pro Arg Arg Glu Gly Ser Phe Pro Gln Gly Gln Leu
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859 Ser Met Leu Gln Glu Lys Ile Asp His Leu Glu Arg Leu Leu Ala Glu
860          50          55          60
864 Asn Asn Glu Ile Ile Ser Asn Ile Arg Asp Ser Val Ile Asn Leu Ser
865 65          70          75          80
868 Glu Ser Val Glu Asp Gly Pro Lys Ser Ser Gln Ser Asn Phe Ser Gln
869          85          90          95
872 Gly Ala Gly Ser His Leu Leu Pro Ser Gln Leu Ser Leu Ser Val Asp
873          100          105          110
877 Thr Ala Asp Cys Leu Phe Ala Ser Gln Ser Gly Ser His Asn Ser Asp
878          115          120          125
881 Val Gln Met Leu Asp Val Tyr Ser Leu Ile Ser Phe Asp Asn Pro Asp
882          130          135          140
886 Gly Gly Val Trp Lys Gln Gly Phe Asp Ile Thr Tyr Glu Ser Asn Glu
887 145          150          155          160
890 Trp Asp Thr Glu Pro Leu Gln Val Phe Val Val Pro His Ser His Asn
891          165          170          175
894 Asp Pro Gly Trp Leu Lys Thr Phe Asn Asp Tyr Phe Arg Asp Lys Thr
895          180          185          190
898 Gln Tyr Ile Phe Asn Asn Met Val Leu Lys Leu Lys Glu Asp Ser Arg
899          195          200          205
902 Arg Lys Phe Ile Trp Ser Glu Ile Ser Tyr Leu Ser Lys Trp Trp Asp
903          210          215          220
906 Ile Ile Asp Ile Gln Lys Lys Asp Ala Val Lys Ser Leu Ile Glu Asn
907 225          230          235          240
910 Gly Gln Leu Glu Ile Val Thr Gly Gly Trp Val Met Pro Asp Glu Ala
911          245          250          255
914 Thr Pro His Tyr Phe Ala Leu Ile Asp Gln Leu Ile Glu Gly His Gln
915          260          265          270
918 Trp Leu Glu Asn Asn Ile Gly Val Lys Pro Arg Ser Gly Trp Ala Ile
919          275          280          285
922 Asp Pro Phe Gly His Ser Pro Thr Met Ala Tyr Leu Leu Asn Arg Ala
923          290          295          300
926 Gly Leu Ser His Met Leu Ile Gln Arg Val His Tyr Ala Val Lys Lys
927 305          310          315          320
930 His Phe Ala Leu His Lys Thr Leu Glu Phe Phe Trp Arg Gln Asn Trp
931          325          330          335
934 Asp Leu Gly Ser Val Thr Asp Ile Leu Cys His Met Met Pro Phe Tyr
935          340          345          350
938 Ser Tyr Asp Ile Pro His Thr Cys Gly Pro Asp Pro Lys Ile Cys Cys
939          355          360          365
942 Gln Phe Asp Phe Lys Arg Leu Pro Gly Gly Arg Phe Gly Cys Pro Trp
943          370          375          380
946 Gly Val Pro Pro Glu Thr Ile His Pro Gly Asn Val Gln Ser Arg Ala
947 385          390          395          400
950 Arg Met Leu Leu Asp Gln Tyr Arg Lys Lys Ser Lys Leu Phe Arg Thr
951          405          410          415

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954 Lys Val Leu Leu Ala Pro Leu Gly Asp Asp Phe Arg Tyr Cys Glu Tyr
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958 Thr Glu Trp Asp Leu Gln Phe Lys Asn Tyr Gln Gln Leu Phe Asp Tyr
959           435           440           445
962 Met Asn Ser Gln Ser Lys Phe Lys Val Lys Ile Gln Phe Gly Thr Leu
963           450           455           460
966 Ser Asp Phe Phe Asp Ala Leu Asp Lys Ala Asp Glu Thr Gln Arg Asp
967 465           470           475           480
970 Lys Gly Gln Ser Met Phe Pro Val Leu Ser Gly Asp Phe Phe Thr Tyr
971           485           490           495
974 Ala Asp Arg Asp Asp His Tyr Trp Ser Gly Tyr Phe Thr Ser Arg Pro
975           500           505           510
978 Phe Tyr Lys Arg Met Asp Arg Ile Met Glu Ser His Leu Arg Ala Ala
979           515           520           525
982 Glu Ile Leu Tyr Tyr Phe Ala Leu Arg Gln Ala His Lys Tyr Lys Ile
983           530           535           540
986 Asn Lys Phe Leu Ser Ser Ser Leu Tyr Thr Ala Leu Thr Glu Ala Arg
987 545           550           555           560
990 Arg Asn Leu Gly Leu Phe Gln His His Asp Ala Ile Thr Gly Thr Ala
991           565           570           575
994 Lys Asp Trp Val Val Val Asp Tyr Gly Thr Arg Leu Phe His Ser Leu
995           580           585           590
998 Met Val Leu Glu Lys Ile Ile Gly Asn Ser Ala Phe Leu Leu Ile Leu
999           595           600           605
1003 Lys Asp Lys Leu Thr Tyr Asp Ser Tyr Ser Pro Asp Thr Phe Leu Glu
1004           610           615           620
1007 Met Asp Leu Lys Gln Lys Ser Gln Asp Ser Leu Pro Gln Lys Asn Ile
1008 625           630           635           640
1011 Ile Arg Leu Ser Ala Glu Pro Arg Tyr Leu Val Val Tyr Asn Pro Leu
1012           645           650           655
1015 Glu Gln Asp Arg Ile Ser Leu Val Ser Val Tyr Val Ser Ser Pro Thr
1016           660           665           670
1019 Val Gln Val Phe Ser Ala Ser Gly Lys Pro Val Glu Val Gln Val Ser
1020           675           680           685
1023 Ala Val Trp Asp Thr Ala Asn Thr Ile Ser Glu Thr Ala Tyr Glu Ile
1024           690           695           700
1027 Ser Phe Arg Ala His Ile Pro Pro Leu Gly Leu Lys Val Tyr Lys Ile
1028 705           710           715           720
1031 Leu Glu Ser Ala Ser Ser Asn Ser His Leu Ala Asp Tyr Val Leu Tyr
1032           725           730           735
1035 Lys Asn Lys Val Glu Asp Ser Gly Ile Phe Thr Ile Lys Asn Met Ile
1036           740           745           750
1039 Asn Thr Glu Glu Gly Ile Thr Leu Glu Asn Ser Phe Val Leu Leu Arg
1040           755           760           765
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1044           770           775           780
1047 Lys His His Glu Val Asn Val Gln Phe Ser Trp Tyr Gly Thr Thr Ile
1048 785           790           795           800
1051 Lys Arg Asp Lys Ser Gly Ala Tyr Leu Phe Leu Pro Asp Gly Asn Ala

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Input Set : A:\SeqList 1975\_0180003.ST25.txt

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1068	865						870				875					880
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1075	Asn	Gly	Tyr	Gln	Ile	Gln	Pro	Arg	Met	Thr	Leu	Ser	Lys	Leu	Pro	Leu
1076				900					905					910		
1079	Gln	Ala	Asn	Val	Tyr	Pro	Met	Thr	Thr	Met	Ala	Tyr	Ile	Gln	Asp	Ala
1080				915				920					925			
1083	Lys	His	Arg	Leu	Thr	Leu	Leu	Ser	Ala	Gln	Ser	Leu	Gly	Val	Ser	Ser
1084				930				935					940			
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1088	945						950				955					960
1091	Asp	Asp	Asn	Arg	Gly	Leu	Glu	Gln	Gly	Ile	Gln	Asp	Asn	Lys	Ile	Thr
1092					965					970						975
1095	Ala	Asn	Leu	Phe	Arg	Ile	Leu	Leu	Glu	Lys	Arg	Ser	Ala	Val	Asn	Thr
1096				980					985					990		
1099	Glu	Glu	Glu	Lys	Lys	Ser	Val	Ser	Tyr	Pro	Ser	Leu	Leu	Ser	His	Ile
1100				995				1000					1005			
1103	Thr	Ser	Ser	Leu	Met	Asn	His	Pro	Val	Ile	Pro	Met	Ala	Asn	Lys	
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1129		1100						1105					1110			
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1133		1115						1120					1125			
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1137		1130						1135					1140			

E--&gt; 1140 Arg ✓

## VERIFICATION SUMMARY

DATE: 01/30/2004

PATENT APPLICATION: US/10/761,435

TIME: 14:54:08

Input Set : A:\SeqList 1975\_0180003.ST25.txt

Output Set: N:\CRF4\01302004\J761435.raw

L:13 M:270 C: Current Application Number differs, Replaced Current Application No

L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date

L:838 M:252 E: No. of Seq. differs, <211> LENGTH:Input:3432 Found:3435 SEQ:17 ✓L:1140 M:252 E: No. of Seq. differs, <211> LENGTH:Input:1143 Found:1144 SEQ:18 ✓